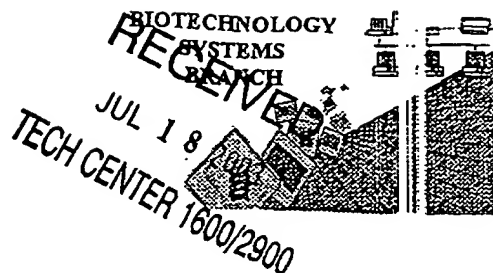


RAW SEQUENCE LISTING
ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/939,226A
Source: 1600
Date Processed by STIC: 7/9/2003

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>>, EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
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Revised 04/24/2003

03 SEP 15 PM 3:17

03 SEP 15 11:38:13



1600

RAW SEQUENCE LISTING

DATE: 07/09/2003

PATENT APPLICATION: US/09/939,226A

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

3 <110> APPLICANT: Euroscreen S.A.
 4 SAMSON, Michel
 5 PARMENTIER, Marc
 6 VASSART, Gilbert
 7 LIBERT, Frederick
 9 <120> TITLE OF INVENTION: Methods for Identifying Compounds which Bind the Active CCR5
 10 Chemokine Receptor
 12 <130> FILE REFERENCE: 9409/2023C
 14 <140> CURRENT APPLICATION NUMBER: 09/939,226A
 15 <141> CURRENT FILING DATE: 2001-08-24
 17 <150> PRIOR APPLICATION NUMBER: US 08/833,752
 18 <151> PRIOR FILING DATE: 1997-04-09
 20 <150> PRIOR APPLICATION NUMBER: US 09/626,939
 21 <151> PRIOR FILING DATE: 2000-07-27
 23 <150> PRIOR APPLICATION NUMBER: US 08/810,028
 24 <151> PRIOR FILING DATE: 1997-03-04
 26 <160> NUMBER OF SEQ ID NOS: 18
 28 <170> SOFTWARE: PatentIn version 3.2
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 31 <211> LENGTH: 792
 32 <212> TYPE: DNA
 33 <213> ORGANISM: Homo sapiens
 35 <400> SEQUENCE: 1
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 38 tcccttcaact acaaaacttc attgcttggc caaaaagaga gttaattcaa tgtagacatc 120
 40 tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact 180
 42 aaatacatc taggacttta taaaagatca ctttttattt atgcacaggg tggaacaaga 240
 44 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc 300
 46 aaaaaatcaa tgtgaagcaa atgcagagccc gcctcctgcc tccgctctac tcaactggtg 360
 48 tcatctttgg ttttgtgggc aacatgctgg tcatcctcat cctgataaac tgcaaaaggc 420
 50 tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg ttttcccttc 480
 52 ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt 540
 54 gtcaactctt gacagggtc tattttatag gcttcttctc tgggaatcttc ttcacatcc 600
 56 tcctgacaat cgataggta ctaggtgtcg tccatgctgt gtttgcttta aaagccagga 660
 58 cggtcacctt tggggtgtg acaagtgtga tcacttgggt ggtggctgtg tttgcgtctc 720
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 66 <211> LENGTH: 1477
 67 <212> TYPE: DNA
 68 <213> ORGANISM: Homo sapiens
 71 <220> FEATURE:
 72 <221> NAME/KEY: misc_feature

Does Not Comply
 Corrected Diskette Needed

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

00SEP15 PM 3:10

SEQUENCE LISTING

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 74 <223> OTHER INFORMATION: Any nucleotide
 76 <220> FEATURE:
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 86 tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact 180
 88 aaatacattc taggacttta taaaagatca ctttttattt atgcacaggg tggacaaga 240
 90 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc 300
 92 aaaaaatcaa tgtgaagcaa atcgagccc gcctcctgcc tccgctctac tcaactggtg 360
 94 tcatcttttg ttttggggc aacatgctgg tcatcctcat cctgataaac tgcaaaaggg 420
 96 tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg tttttccttc 480
 98 ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt 540
 100 gtcaactctt gacagggctc ttttttatag gcttcttctc tggaaatctc tcatcatcc 600
 102 tccgtgacaat cgataggtag ctggctgtcg tccatgctgt gtttgcttta aaagccagga 660
 104 cggtcacctt tggggtggtg acaagtgtga tcaactgggt ggtggctgtg tttgcgtctc 720
 106 tcccaggaat catctttacc agatctcaaa aagaagggtc tcattacacc tgcagctctc 780
 108 attttccata cagtcagtat caattctgga agaatttcca gacattaaag atagtcatct 840
 110 tggggctggt cctgccgctg cttgtcatgg tcatctgcta ctcggaatc ctaaaaaactc 900
 112 tgcttcggtg tcgaaatgag aagaagaggc acagggctgt gaggcttate ttcaccatca 960
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 116 aggaattctt tggcctgaat aattgcagta gctctaacag gttggaccaa gctatgcagg 1080
 118 tgacagagac tottgggatg acgcactgct gcataaccc catcatctat gcctttgtcg 1140
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 126 acccagtcag agttgtgcac atggcttagt tttcatacac agcctgggct ggggggtnggt 1380
 128 tggngaggt cttttttaaa aggaagttac tgttatagag ggtctaagat tcatccattt 1440
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135 <212> TYPE: DNA

136 <213> ORGANISM: Homo sapiens

138 <400> SEQUENCE: 3

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 141 tcccttcaact acaaaacttc attgcttggc caaaaagaga gtttaattcaa tgtagacatc 120
 143 tatgtaggca attaaaaacc tattgatgta taaaacagtt tgcattcatg gagggcaact 180
 145 aaatacattc taggacttta taaaagatca ctttttattt atgcacaggg tggacaaga 240
 147 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc 300
 149 aaaaaatcaa tgtgaagcaa atcgagccc gcctcctgcc tccgctctac tcaactggtg 360
 151 tcatcttttg ttttggggc aacatgctgg tcatcctcat cctgataaac tgcaaaaggg 420
 153 tgaagagcat gactgacatc tacctgctca acctggccat ctctgacctg tttttccttc 480
 155 ttactgtccc cttctgggct cactatgctg ccgcccagtg ggactttgga aatacaatgt 540
 157 gtcaactctt gacagggctc ttttttatag gcttcttctc tggaaatctc tcatcatcc 600
 159 tccgtgacaat cgataggtag ctggctgtcg tccatgctgt gtttgcttta aaagccagga 660
 161 cggtcacctt tggggtggtg acaagtgtga tcaactgggt ggtggctgtg tttgcgtctc 720

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

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163 tcccaggaat catctttacc agatctcaaa aagaaggctct tcattacacc tgcagctctc 780
165 attttccata cattaaagat agtcatcttg gggctgggtcc tgccgctgct tgtcatgggc 840
167 atctgctact cggaatcct aaaaactctg ctctgggtgc gaaatgagaa gaagaggcac 900
169 agggctgtga ggcttatctt caccatcatg attgtttatt ttctctcttg ggctccctac 960
171 aacattgtcc ttctcctgaa caccttccag gaattctttg gcctgaataa ttgcagtagc 1020
173 tctaacaggt tggaccaagc tatgcagggt acagagactc ttgggatgac gcactgctgc 1080
175 atcaacccca tcatctatgc ctttgcggg gagaagttca gaaactacct cttagtcttc 1140
177 ttccaaaagc acattgccaa acgcttctgc aaatgctgtt ctattttcca gcaagaggct 1200
179 cccgagcgag caagctcagt ttacaccgca tccactgggg agcaggaaat atctgtgggc 1260
181 ttgtgacacg gactcaagtg ggctgggtgac ccagtcagag ttgtgcacat ggcttagttt 1320
183 tcatacacag cctgggctgg ggggtggttg gaggtctttt ttaaaaggaa gttactgtta 1380
185 tagagggtct aagattcatt ctttatttg gcatctgttt aaagtagatt agatccgaat 1440
187 tc 1442
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191 <211> LENGTH: 184
192 <212> TYPE: PRT
193 <213> ORGANISM: Homo sapiens
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198 1 5 10 15
201 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
202 20 25 30
205 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
206 35 40 45
209 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
210 50 55 60
213 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
214 65 70 75 80
217 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
218 85 90 95
221 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
222 100 105 110
225 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
226 115 120 125
229 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
230 130 135 140
233 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
234 145 150 155 160
237 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
238 165 170 175
241 Thr Cys Ser Ser His Phe Pro Tyr
242 180
245 <210> SEQ ID NO: 5
246 <211> LENGTH: 352
247 <212> TYPE: PRT
248 <213> ORGANISM: Homo sapiens
250 <400> SEQUENCE: 5
252 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
253 1 5 10 15

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

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256 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
257          20          25          30
260 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
261          35          40          45
264 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
265          50          55          60
268 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
269 65          70          75          80
272 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
273          85          90          95
276 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
277          100          105          110
280 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
281          115          120          125
284 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
285          130          135          140
288 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
289 145          150          155          160
292 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
293          165          170          175
296 Thr Cys Ser Ser His Phe Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn
297          180          185          190
300 Phe Gln Thr Leu Lys Ile Val Ile Leu Gly Leu Val Leu Pro Leu Leu
301          195          200          205
304 Val Met Val Ile Cys Tyr Ser Gly Ile Leu Lys Thr Leu Leu Arg Cys
305          210          215          220
308 Arg Asn Glu Lys Lys Arg His Arg Ala Val Arg Leu Ile Phe Thr Ile
309 225          230          235          240
312 Met Ile Val Tyr Phe Leu Phe Trp Ala Pro Tyr Asn Ile Val Leu Leu
313          245          250          255
316 Leu Asn Thr Phe Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser
317          260          265          270
320 Asn Arg Leu Asp Gln Ala Met Gln Val Thr Glu Thr Leu Gly Met Thr
321          275          280          285
324 His Cys Cys Ile Asn Pro Ile Ile Tyr Ala Phe Val Gly Glu Lys Phe
325          290          295          300
328 Arg Asn Tyr Leu Leu Val Phe Phe Gln Lys His Ile Ala Lys Arg Phe
329 305          310          315          320
332 Cys Lys Cys Cys Ser Ile Phe Gln Gln Glu Ala Pro Glu Arg Ala Ser
333          325          330          335
336 Ser Val Tyr Thr Arg Ser Thr Gly Glu Gln Glu Ile Ser Val Gly Leu
337          340          345          350
340 <210> SEQ ID NO: 6
341 <211> LENGTH: 215
342 <212> TYPE: PRT
343 <213> ORGANISM: Homo sapiens
345 <400> SEQUENCE: 6
347 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
348 1          5          10          15

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

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351 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
352      20      25      30
355 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
356      35      40      45
359 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
360      50      55      60
363 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
364 65      70      75      80
367 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
368      85      90      95
371 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
372      100     105     110
375 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
376      115     120     125
379 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
380      130     135     140
383 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
384 145     150     155     160
387 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
388      165     170     175
391 Thr Cys Ser Ser His Phe Pro Tyr Ile Lys Asp Ser His Leu Gly Ala
392      180     185     190
395 Gly Pro Ala Ala Ala Cys His Gly His Leu Leu Leu Gly Asn Pro Lys
396      195     200     205
399 Asn Ser Ala Ser Val Ser Lys
400      210     215
403 <210> SEQ ID NO: 7
404 <211> LENGTH: 360
405 <212> TYPE: PRT
406 <213> ORGANISM: Homo sapiens
409 <220> FEATURE:
410 <221> NAME/KEY: MISC_FEATURE
411 <222> LOCATION: (325)..(327)
412 <223> OTHER INFORMATION: Xaa = any amino acid
414 <400> SEQUENCE: 7
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417 1      5      10      15
420 Gly Glu Glu Val Thr Thr Phe Phe Asp Tyr Asp Tyr Gly Ala Pro Cys
421      20      25      30
424 His Lys Phe Asp Val Lys Gln Ile Gly Ala Gln Leu Leu Pro Pro Leu
425      35      40      45
428 Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn Met Leu Val Val
429      50      55      60
432 Leu Ile Leu Ile Asn Cys Lys Lys Leu Lys Cys Leu Thr Asp Ile Tyr
433 65      70      75      80
436 Leu Leu Asn Leu Ala Ile Ser Asp Leu Leu Phe Ile Ile Thr Leu Pro
437      85      90      95
440 Leu Trp Ala His Ser Ala Ala Asn Glu Trp Val Phe Gly Asn Ala Met
441      100     105     110

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09/939,226A 15

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<213> ORGANISM: Artificial Sequence
<220> FEATURE:
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see p.7 for error explanation...

17

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<213> ORGANISM: Artificial Sequence
<220> FEATURE:
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18

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<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: :
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ctgatctaga gccatgtgca caactct

21

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003
TIME: 15:53:55

FYI

Input Set : A:\sequencelisting.txt
Output Set: N:\CRF4\07092003\I939226A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <2:()> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; N Pos. 1377,1384,1385
Seq#:7; Xaa Pos. 325,326,327
Seq#:8; Xaa Pos. 231,232,233,333,334,335
Seq#:10; Xaa Pos. 145,146,147,321,322,323

✓ OK

Use of <220> Feature(NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

Seq#:14,15,16,17

↑
error explanation

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003

TIME: 15:53:55

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1320
M 341 Repeated in SeqNo=2
L 496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:320
L 582 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:224
M 341 Repeated in SeqNo=8
L:771 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:144
M:341 Repeated in SeqNo=10
L:885 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:887 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:14, <213>
ORGANISM:Artificial Sequence
L:887 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:14, <213>
ORGANISM:Artificial Sequence
L:887 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:887
L:894 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:896 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:15, <213>
ORGANISM:Artificial Sequence
L:896 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:15, <213>
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L:896 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:15,Line#:896
L:903 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:905 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:16, <213>
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L:912 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
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ORGANISM:Artificial Sequence
L:914 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:17, <213>
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L:914 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:17,Line#:914